

SEQUENCE LISTING

<110> DeFrenne, Catherine
DelMelle, Christine
Ruelle, Jean-Louis

<120> Novel Compounds

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<212> DNA

<213> Neisseria meningitidis

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cgc	ccgcgtt	cgactttcg	ggcaactgg	atttcacg	gcaacacaaa	1500
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tcc	tcgaactcgc	gtgggctac	gaaggcgacc	gctggcaata	caatctggca	1680
ctc	accgctcgg	caactacatt	tacgccc	ccttaa	cggacgcggc	1740
ccc	tcgaagacga	cagcggaaatg	aagctcgtgc	gctacaacca	atccgg	1800
gac	gcgcggaa	cggaaatctac	ttcaaacg	caccgcgc	ccgcac	1860
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gct	tcggcttcc	cctgaaagcc	tcgctgacc	accgcata	tgccaattt	2040
gact	gcgtgttgc	ccaaaacaaa	ctcgccgct	acgaaacg	cacccc	2100
cac	tcaacctcgg	cggaaactac	cggcata	cgcgtat	cgagtgg	2160
tgg	tacgtca	aagccacaa	cctgctcaac	caatcg	ccacag	2220
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<211> 758

<212> PRT

<213> Neisseria meningitidis

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Leu	Glu	Thr	Val	Thr	Val	Val	Gly	Lys	Ser	Arg	Pro	Arg	Ala	Thr	Ser
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Gly	Leu	Leu	His	Thr	Ser	Thr	Ala	Ser	Asp	Lys	Ile	Ile	Ser	Gly	Asp
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Thr	Leu	Arg	Gln	Lys	Ala	Val	Asn	Leu	Gly	Asp	Ala	Leu	Asp	Gly	Val
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Pro	Gly	Ile	His	Ala	Ser	Gln	Tyr	Gly	Gly	Gly	Ala	Ser	Ala	Pro	Val
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Ile	Arg	Gly	Gln	Thr	Gly	Arg	Arg	Ile	Lys	Val	Leu	Asn	His	His	Gly
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Glu	Thr	Gly	Asp	Met	Ala	Asp	Phe	Ser	Pro	Asp	His	Ala	Ile	Met	Val
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Asp	Thr	Ala	Leu	Ser	Gln	Gln	Val	Glu	Ile	Leu	Arg	Gly	Pro	Val	Thr
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Leu	Leu	Tyr	Ser	Ser	Gly	Asn	Val	Ala	Gly	Leu	Val	Asp	Val	Ala	Asp
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Gly	Lys	Ile	Pro	Glu	Lys	Met	Pro	Glu	Asn	Gly	Val	Ser	Gly	Glu	Leu
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Ile	Asn	Ile	Gly	Leu	Gly	Lys	Asn	Phe	Val	Leu	His	Thr	Glu	Gly	Leu
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															200
															205

Tyr Arg Lys Ser Gly Asp Tyr Ala Val Pro Arg Tyr Arg Asn Leu Lys
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 Arg Leu Pro Asp Ser Pro Arg Arg Phe Ala Asn Gly Gln His Arg Ala
 225 230 235 240
 Val Leu Gly Trp Arg Lys Arg Phe Tyr Arg Arg Thr Tyr Ser Asp Arg
 245 250 255
 Arg Asp Gln Tyr Gly Leu Pro Ala His Ser His Glu Tyr Asp Asp Cys
 260 265 270
 His Ala Asp Ile Ile Trp Gln Lys Ser Leu Ile Asn Lys Arg Tyr Leu
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 Gln Leu Tyr Pro His Leu Leu Thr Glu Glu Asp Val Asp Tyr Asp Asn
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 Ala His Asn Gly Lys Pro Trp Ile Asp Leu Arg Asn Lys Arg Tyr Glu
 325 330 335
 Leu Arg Ala Glu Trp Lys Gln Pro Phe Pro Gly Phe Glu Ala Leu Arg
 340 345 350
 Val His Leu Asn Arg Asn Asp Tyr His His Asp Glu Lys Ala Gly Asp
 355 360 365
 Ala Val Glu Asn Phe Phe Asn Asn Gln Thr Gln Asn Ala Arg Ile Glu
 370 375 380
 Leu Arg His Gln Pro Ile Gly Arg Leu Lys Gly Ser Trp Gly Val Gln
 385 390 395 400
 Tyr Leu Gly Gln Lys Ser Ser Ala Leu Ser Ala Thr Ser Glu Ala Val
 405 410 415
 Lys Gln Pro Met Leu Leu Asp Asn Lys Val Gln His Tyr Ser Phe Phe
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 Gly Val Glu Gln Ala Asn Trp Asp Asn Phe Thr Leu Glu Gly Val
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 Arg Val Glu Lys Gln Lys Ala Ser Ile Arg Tyr Asp Lys Ala Leu Ile
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 Asp Arg Glu Asn Tyr Tyr Lys Gln Pro Leu Pro Asp Leu Gly Ala His
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 Arg Gln Thr Ala Arg Ser Phe Ala Leu Ser Gly Asn Trp Tyr Phe Thr
 485 490 495
 Pro Gln His Lys Leu Ser Leu Thr Ala Ser His Gln Glu Arg Leu Pro
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 Ser Thr Gln Glu Leu Tyr Ala His Gly Lys His Val Ala Thr Asn Thr
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 Phe Glu Val Gly Asn Lys His Leu Asn Lys Glu Arg Ser Asn Asn Ile
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 Glu Leu Ala Leu Gly Tyr Glu Gly Asp Arg Trp Gln Tyr Asn Leu Ala
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 Leu Tyr Arg Asn Arg Phe Gly Asn Tyr Ile Tyr Ala Gln Thr Leu Asn
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 Asp Gly Arg Gly Pro Lys Ser Ile Glu Asp Asp Ser Glu Met Lys Leu
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 Val Arg Tyr Asn Gln Ser Gly Ala Asp Phe Tyr Gly Ala Glu Gly Glu
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 Ile Tyr Phe Lys Pro Thr Pro Arg Tyr Arg Ile Gly Val Ser Gly Asp
 610 615 620
 Tyr Val Arg Gly Arg Leu Lys Asn Leu Pro Ser Leu Pro Gly Arg Glu
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 Asp Ala Tyr Gly Asn Arg Pro Phe Ile Ala Gln Asp Asp Gln Asn Ala
 645 650 655
 Pro Arg Val Pro Ala Ala Arg Leu Gly Phe His Leu Lys Ala Ser Leu

660	665	670
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675	680	685
Asn Lys Leu Ala Arg Tyr Glu Thr Arg Thr Pro Gly His His Met Leu		
690	695	700
Asn Leu Gly Ala Asn Tyr Arg Arg Asn Thr Arg Tyr Gly Glu Trp Asn		
705	710	715
Trp Tyr Val Lys Ala Asp Asn Leu Leu Asn Gln Ser Val Tyr Ala His		
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<212> DNA

<213> Neisseria meningitidis

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gttaccgc acggacagct catcaaagaa acgcccacaa ccatcgatac gctcaatatac	240
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<212> PRT
<213> Neisseria meningitidis

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35 40 45
Val Leu Lys Gly Tyr Ile Asn Tyr Asp Glu Ala Ala Val Thr Arg Asn
50 55 60
Gly Gln Leu Ile Lys Glu Thr Pro Gln Thr Ile Asp Thr Leu Asn Ile
65 70 75 80
Gln Lys Asn Lys Asn Tyr Gly Thr Asn Asp Leu Ser Ser Ile Leu Glu
85 90 95
Gly Asn Ala Gly Ile Asp Ala Ala Tyr Asp Met Arg Gly Glu Ser Ile
100 105 110
Phe Leu Arg Gly Phe Gln Ala Asp Ala Ser Asp Ile Tyr Arg Asp Gly
115 120 125
Val Arg Glu Ser Gly Gln Val Arg Arg Ser Thr Ala Asn Ile Glu Arg
130 135 140
Val Glu Ile Leu Lys Gly Pro Ser Ser Val Leu Tyr Gly Arg Thr Asn
145 150 155 160
Gly Gly Gly Val Ile Asn Met Val Ser Lys Tyr Ala Asn Phe Lys Gln
165 170 175
Ser Arg Asn Ile Gly Ala Val Tyr Gly Ser Trp Ala Asn Arg Ser Leu
180 185 190
Asn Met Asp Ile Asn Glu Val Leu Asn Lys Asn Val Ala Ile Arg Leu
195 200 205
Thr Gly Glu Val Gly Arg Ala Asn Ser Phe Arg Ser Gly Ile Asp Ser
210 215 220
Lys Asn Val Met Val Ser Pro Ser Ile Thr Val Lys Leu Asp Asn Gly
225 230 235 240
Leu Lys Trp Thr Gly Gln Tyr Thr Tyr Asp Asn Val Glu Arg Thr Pro
245 250 255
Asp Arg Ser Pro Thr Lys Ser Val Tyr Asp Arg Phe Gly Leu Pro Tyr
260 265 270
Arg Met Gly Phe Ala His Arg Asn Asp Phe Val Lys Asp Lys Leu Gln
275 280 285
Val Trp Arg Ser Asp Leu Glu Tyr Ala Phe Asn Asp Lys Trp Arg Ala
290 295 300
Gln Trp Gln Leu Ala His Arg Thr Ala Ala Gln Asp Phe Asp His Phe
305 310 315 320
Tyr Ala Gly Ser Glu Asn Gly Asn Leu Ile Lys Arg Asn Tyr Ala Trp
325 330 335
Gln Gln Thr Asp Asn Lys Thr Leu Ser Ser Asn Phe Thr Leu Asn Gly
340 345 350
Asp Tyr Thr Ile Gly Arg Phe Glu Asn His Leu Thr Val Gly Met Asp
355 360 365
Tyr Ser Arg Glu His Arg Asn Pro Thr Leu Gly Phe Arg Arg Asn Phe
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Thr Ala Ser Ile Asp Pro Tyr Asp Arg Ala Ser Arg Pro Ala Ser Gly
385 390 395 400
Arg Leu Gln Arg Ile Leu Ala Gln Asp Arg His Lys Ala Asp Ser Tyr

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465	470	475
Tyr Asn Lys Ala Phe Ala Pro Tyr Gly Gly Arg Gly Gly Tyr Leu Ser		
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Ile Asn Thr Ser Ser Ala Val Phe Asn Ala Asp Pro Glu Tyr Thr		
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Ser Thr Thr Leu Ser Ala Tyr Gln Ile Glu Arg Phe Asn Ile Arg Tyr		
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545	550	555
His Arg Ser Arg Gly Val Glu Leu Ser Ala Ile Gly Gln Ile Ile Pro		
565	570	575
Lys Lys Leu Tyr Leu Arg Gly Ser Leu Gly Val Met Gln Ala Lys Val		
580	585	590
Val Glu Asp Lys Lys Asn Pro Asp Arg Val Gly Ile His Leu Asn Asn		
595	600	605
Thr Ser Asn Val Thr Gly Asn Leu Phe Phe Arg Tyr Thr Pro Thr Glu		
610	615	620
Asn Leu Tyr Gly Glu Ile Gly Val Thr Gly Thr Gly Lys Arg Tyr Gly		
625	630	635
Tyr Asn Ser Arg Asn Lys Glu Val Thr Thr Leu Pro Gly Phe Ala Arg		
645	650	655
Val Asp Ala Met Leu Gly Trp Asn His Lys Asn Val Asn Val Thr Phe		
660	665	670
Ala Ala Ala Asn Leu Phe Asn Gln Lys Tyr Trp Arg Ser Asp Ser Met		
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<211> 378

<212> DNA

<213> Neisseria meningitidis

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tcccgcaacg gcatcatcaa agaacgcagc aatctgaccg tctatatttga aaacggcgta	300
ctcgccgca ccgaaggcga cgtccctgcaa aacgctgccg aagcgctcaa agaccgcag	360
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<212> PRT

<213> Neisseria meningitidis

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35 40 45
Leu Gly Met Thr Lys Asp Gln Val Leu Leu Leu Leu Gly Ser Pro Leu
50 55 60
Leu Arg Asp Ala Phe His Thr Glu Arg Trp Asp Tyr Thr Phe Asn Thr
65 70 75 80
Ser Arg Asn Gly Ile Ile Lys Glu Arg Ser Asn Leu Thr Val Tyr Phe
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Glu Asn Gly Val Leu Val Arg Thr Glu Gly Asp Val Leu Gln Asn Ala
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<210> 7

<211> 864

<212> DNA

<213> *Neisseria meningitidis*

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<210> 8

<211> 287

<212> PRT

<213> *Neisseria meningitidis*

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Ser	Ala	Ala	Ala	Asp	Asn	Gly	Ala	Glu	Lys	Lys	Glu	Ile	Val	Phe	Gly
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Thr	Thr	Val	Gly	Asp	Phe	Gly	Asp	Met	Val	Lys	Glu	Gln	Ile	Gln	Ala
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Glu	Leu	Glu	Lys	Lys	Gly	Tyr	Thr	Val	Lys	Leu	Val	Glu	Phe	Thr	Asp
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Tyr	Val	Arg	Pro	Asn	Leu	Ala	Leu	Ala	Glu	Gly	Glu	Leu	Asp	Ile	Asn

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Leu Asp Ile Thr Glu Val Phe Gln Val Pro Thr Ala Pro Leu Gly Leu			
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Tyr Pro Gly Lys Leu Lys Ser Leu Glu Glu Val Lys Asp Gly Ser Thr			
130	135	140	
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Leu Asp Glu Leu Gly Trp Ile Lys Leu Lys Asp Gly Ile Asn Pro Leu			
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Thr Ala Ser Lys Ala Asp Ile Ala Glu Asn Leu Lys Asn Ile Lys Ile			
180	185	190	
Val Glu Leu Glu Ala Ala Gln Leu Pro Arg Ser Arg Ala Asp Val Asp			
195	200	205	
Phe Ala Val Val Asn Gly Asn Tyr Ala Ile Ser Ser Gly Met Lys Leu			
210	215	220	
Thr Glu Ala Leu Phe Gln Glu Pro Ser Phe Ala Tyr Val Asn Trp Ser			
225	230	235	240
Ala Val Lys Thr Ala Asp Lys Asp Ser Gln Trp Leu Lys Asp Val Thr			
245	250	255	
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<213> Neisseria meningitidis

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<211> 321

<212> PRT

<213> Neisseria meningitidis

<400> 10

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Ser	Ala	Ala	Thr	Leu	Thr	Val	Pro	Thr	Ala	Arg	Gly	Asp	Ala	Val	Val
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					50			55			60				
Thr	Leu	Thr	Glu	Leu	Gly	Val	Asn	Val	Gly	Ala	Thr	Thr	Ala	Pro	Met
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Arg	Val	Asp	Tyr	Leu	Gln	Pro	Ala	Phe	Asp	Lys	Ala	Ala	Thr	Val	Gly
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Thr	Leu	Phe	Glu	Pro	Asp	Tyr	Glu	Ala	Leu	His	Arg	Tyr	Asn	Pro	Gln
					100			105			110				
Leu	Val	Ile	Thr	Gly	Gly	Pro	Gly	Ala	Glu	Ala	Tyr	Glu	Gln	Leu	Ala
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					130			135			140				
Thr	Ser	Gly	Glu	Lys	Gln	Met	Glu	Thr	Leu	Ala	Arg	Ile	Phe	Gly	Lys
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Lys	Asn	Pro	Asp	Trp	Ile	Phe	Ile	Ile	Asp	Arg	Thr	Ala	Ala	Ile	Gly
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Gln	Glu	Gly	Pro	Ala	Ala	Val	Glu	Val	Leu	Asp	Asn	Ala	Leu	Val	Arg
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Gln	Leu	Lys	Ala	Ala	Phe	Glu	Lys	Ala	Glu	Pro	Val	Ala	Ala	Gly	Lys
					305			310			315			320	
Glu															